# Map of pRM1

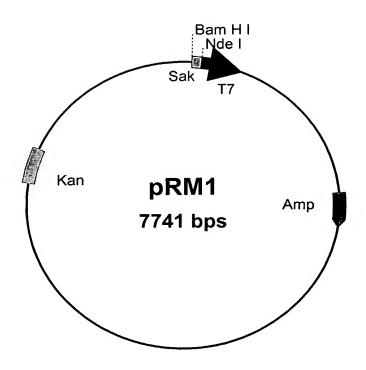


Fig. 1

#### Fig. 2 Sequence of oligonucleotide to construct the protein expression cassette

- 1. 5' GATCAAGCTTATCATCGATAAGCTTACAGGACGCTGGGTTAAAA
  GTATTT -3" (51 mer ) PEC 1
- 2. 5"-

5

10

15

- ATCTTATTGACCTCTCAAAACTTAATCCACATCAAAACTCAAATAC
  TTTTAACCC -3" (55 mer ) PEC 2
- 3. 5"-AGAGGTCAATAAGATTATAATATGTGATGCTTCACAATTCTGATG
  TATGGCAAAA -3" (55 mer) PEC 3
- 4. 5"-ATGAGGTCTTCCTTAAGTTCATTATTGGTTTTTGCCATACATCA GAATT -3" (50 mer) PEC 4

# Fig. 3 Complete nucleotide sequence of expression cassette, OXY-1 GATCAAGCTTATCATCGATAAGCTTACAGGACGCTGGGTTAAAAGTATTTG AGTTTTGATGTGGATTAAGTTTTGAGAGGTCAATAAGATTATAATATGTGA TGCTTCACAATTCTGATGTATGGCAAAACCATAATAATGAACTTAAGGAAG ACCTCATG (161 mer )

#### Fig.4 Nucleotide sequence of SAK gene

30

(411 nucleotides)

#### Fig.5 Nucleotide sequence of SAK 1 gene

15 Oligo's:

SAK-3 primer:

5'- GAACTTAAGGAAGATATACATATGTCAAGTTCATTCGACAAAGGA-3' (45 mer )

SAK-2 primer:

20 5'- CGGCTACTGGATCCTCCACTTTTATCCAAACTGATTT -3' (38 mer)

## √ <sup>th</sup> 'U 015118-6

#### Fig.6 Nucleotide sequence of SAK 2 gene

15 SAK-4 primer :

5'- GAACTTAAGCATATGGCTGGAGCTTATAAAAAGGGC-3' ( 36 mer) SAK-2 primer:

2. 5'- CGGCTACTGGATCCTCCACTTTTATCCAAACTGATTT -3' ( 38 mer

20

25

30

Fig. 7 Schematic representation of construction of plasmid, pOXYPRO, carrying OXY-1 expression cassette.

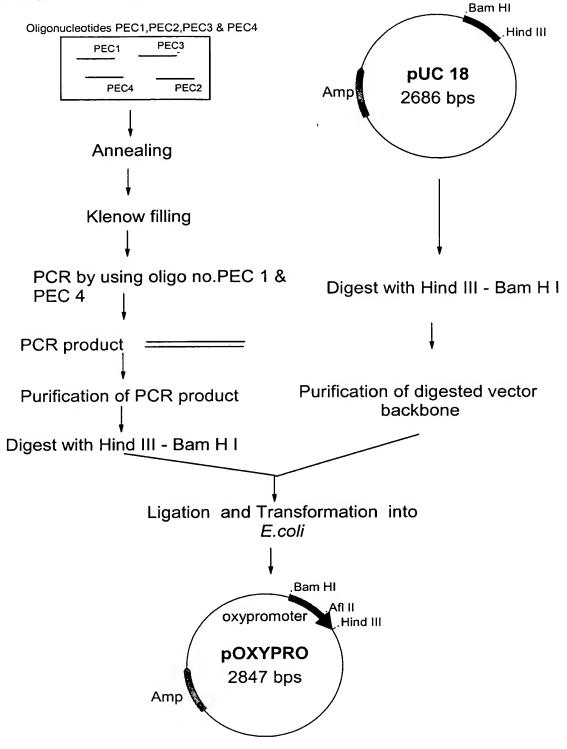


Fig. 8 Construction and map of plasmid, pOXYSAK-1 containing SAK-1 gene under OXY-1 expression cassette.

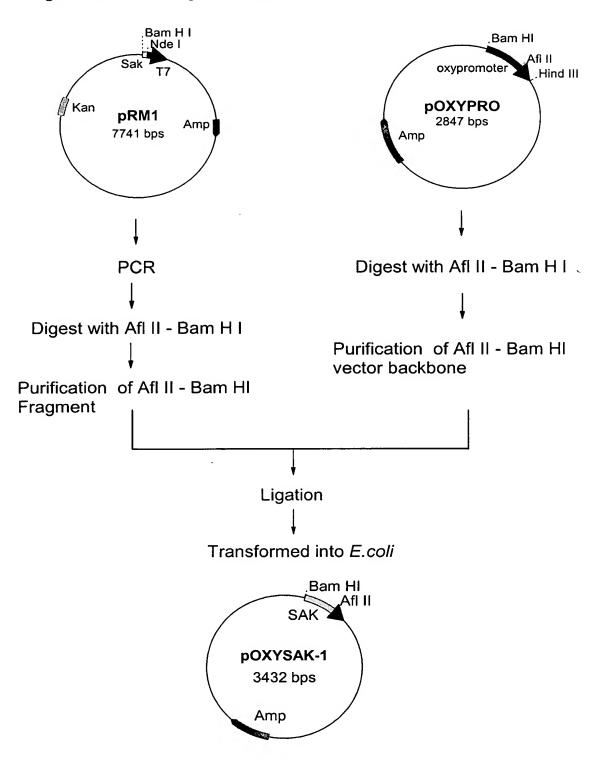


Fig.9 Construction and map of plasmid, pOXYSAK-2 containing SAK-2 gene under OXY-1 expression cassette.

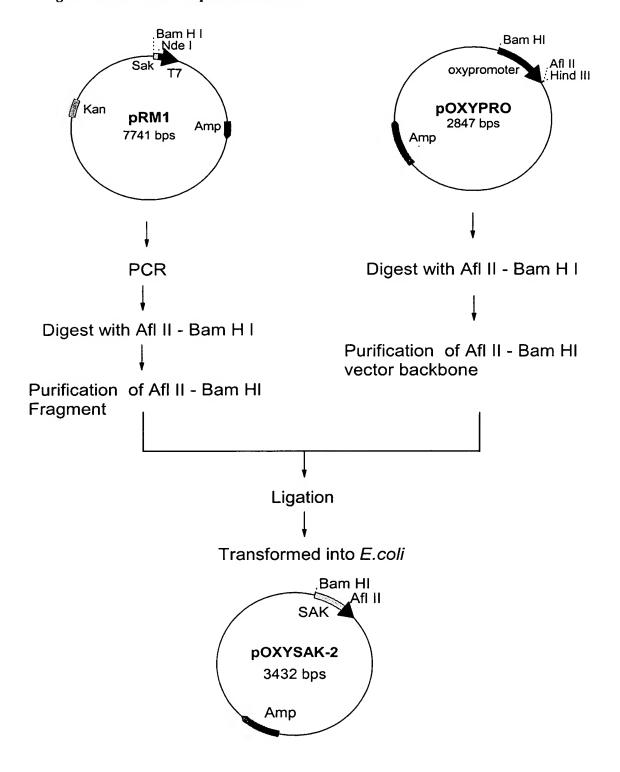


Fig.10 Sequences of SAK, SAK-1 and SAK-2 genes.

# **⋖ 👫 ″U 015118-6**

Terminator sequence in SAK-1 and SAK-2 are shown in bold letters and modified residues in SAK-2 are shown is small bold letters.

5	S	A	K

TCAAGTTCATTCGACAAAGGAAA

SAK-1

GAACTTAAGGAAGATATACATATGTCAAGTTCATTCGACAAAGGAAA
SAK-2
GAACTTAAGCATATG g c tGGA

10 gc

SAK

ATATAAAAAGGCCGATGACCCAGTTATTTTGAACCAACAGGCCCGT SAK-1

15 ATATAAAAAGGGCGATGACGCGAGTTATTTTGAACCAACAGGCCCGT SAK-2

t

TATAAAAAGGGCGATGACGCGAGTTATTTTGAACCAACAGGCCCGT

SAK

20 ATTTGATGGTAAATGTGACTGGAGTTGATGGTAAAGGAAATGAATTG
SAK-1
ATTTGATGGTAAATGTGACTGGAGTTGATGGTAAAGGAAATGAATTG
SAK-2

ATTTGATGGTAAATGTGACTGGAGTTGATGGTAAAGGAAATGAATTG

25

SAK CTATCCCCTCATTA TGTCGAGTTTCCTATTAAACCTGGGACTACACT SAK-1 CTATCCCCTCATTA TGTCGAGTTTCCTATTAAACCTGGGACTACACT SAK-2 CTATCCCCTCATTA TGTCGAGTTTCCTATTAAACCTGGGACTACACT

30 SAK

TACAAAAGAAAAATTGAATACTATGTCGAATGGGCATTAGATGCGA SAK-1

TACAAAAGAAAAATTGAATACTATGTCGAATGGGCATTAGATGCGA

	SAK-2
	TACAAAAGAAAAATTGAATACTATGTCGAATGGGCATTAGATGCGA
	SAK
5	CAGCATATAAAGAGTTTAGAGTAGTTGAATTAGATCCAAGCGCAAAG
	SAK-1
	CAGCATATAAAGAGTTTAGAGTTGAATTAGATCCAAGCGCAAAG
	SAK-2
	CAGCATATAAAGAGTTTAGAGTAGTTGAATTAGATCCAAGCGCAAAG
10	
	SAK
	ATCGAAGTCACTTATTATGATAAGAATAAGAAAAAAAGAAGAAACGAA
15	SAK-1
	ATCGAAGTCACTTATTATGATAAGAATAAGAAAAAAGAAGAAACGAA
	SAK-2
	ATCGAAGTCACTTATTATGATAAGAATAAGAAAAAAAAGAAGAAACGAA
20	
	SAK
	GTCTTTCCCTATAACAGAAAAAGGTTTTGTTGTCCCAGATTTATCAGA
	SAK-1
25	GTCTTTCCCTATAACAGAAAAAGGTTTTGTTGTCCCAGATTTATCAGA
	SAK-2
	GTCTTTCCCTATAACAGAAAAAGGTTTTGTTGTCCCAGATTTATCAGA
	SAK GCATATTAAAAACCCTGGATTCAACTTAATTACAAAGGTTGTTATAG
30	SAK-1
50	GCATATTAAAAACCCTGGATTCAACTTAATTACAAAGGTTGTTATAG
	SAK-2
	GCATATTAAAAACCCTGGATTCAACTTAATTACAAAGGTTGTTATAG

SAK AAAAGAAATAA

	SAK-1						
	AAAAGAAATA	AAACAAAAT	<b>AGTTGTTTA</b>	TTATAGAAAG	STAATGTC		
	SAK-2						
5	AAAAGAAATA	AAACAAAAT	AGTTGTTTA	TTATAGAAAG	TAATGTC		
	SAK-1						
	TTGATTGAAT	ATGTGTAGT	GAAATTATC	TTTCATCAAA	TTCTCATT		
	SAK-2						
10	TTGATTGAAT	ATGTGTAGT	GAAATTATC	TTTCATCAAA	TTCTCATT		
	SAK-1						
	CATGCACGAA	ATGGTTCTG	CCCCACCTAA	TCAGATATTA	ACGTGACT		
	SAK-2						
15	CATGCACGAA	ATGGTTCTGC	CCCCACCTAA	TCAGATATTA	ACGTGACT		
	SAK-1						
		• • • <del>• • • • • • • • • • • • • • • • </del>		CTCCACCATC	CACTACCC		
20	TATGGGGAGAAATCAGTTTGGATAAAAGTGGAGGATCCAGTAGCC SAK-2						
20	TATGGGGAGA	A A A T C A G T T 1	ГССАТАААА	CTGGAGGATO	CAGTAGCC		
	TATOGGAG						
	SAK-1 <b>G</b>						
25	SAK-2 <b>G</b>						
	Fig. 11 Modification	ation of SAK i	in SAK-2 . Mo	dified residues	are shown in b	old	
	letters.						
	. 1	10	20	30	40		
30	SAK						
	SSSFD <b>K</b> G <b>K</b> TK	KGDDASYFEP	TGPYLMVNVI	GVDGKGNELI	SPHYVEFP		
	SAK-2						
	<b>A</b> G <b>A</b> TKKGDDA	SYFEPTGPYL	MVNVTGVDG	KGNELLSPHY	VEFP		

## \*U 015118-6

SAK IKPGTTLTKEKIEYYVEWALDATAYKEFRVVELAPSAKIEVTYYDKNKK SAK-2 IKPGTTLTKEKIEYYVEWALDATAYKEFRVVELAPSAKIEVTYYDKNKK SAK EETTKSFPITEKGFVVPDLSEHIKNPGFNLITKVVIEKK SAK-2 EETTKSFPITEKGFVVPDLSEHIKNPGFNLITKVVIEKK